11-1272

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208767

ACCESS DB #_______PLEASE PRINT CLEARLY

Scientific and Technical Information Center

SEARCH REQUEST FORM

Requester's Full Name: $\pm ANC 2ARA$ E	xaminer #: 77514 Date: 11-27-06
Art Unit: 163 Phone Number: 2- 576	Serial Number: 10/077.11/
Location (Bldg/Room#): 1459 (Mailbox #): 2018 Res	
***************	*********
To ensure an efficient and quality search, please attach a copy of the cover s	sheet, claims, and abstract or fill out the follov
Title of Invention: ID & Cloning of	Ret 16
Inventors (please provide full names):	detol.
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i /	
Earliest Priority Date: 2/15/02	^
Search Topic:	
Please provide a detailed statement of the search topic, and describe as specific elected species or structures, keywords, synonyms, acronyms, and registry nun Define any terms that may have a special meaning. Give examples or relevant	nbers, and combine with the concept or utility of the invention.
*For Sequence Searches Ont ** Please include all pertinent information (pare	ent. child. divisional. or issued natent numbers) along with the
appropriate serial number.	min, committee and a second parents in an action along main the
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GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd
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OM nucleic - nucleic search, using sw model

December 3, 2006, 02:54:58; Search time 784 Seconds (without alignments) 9195.652 Million cell updates/sec Run on:

Title: Perfect score: Sequence:

US-10-077-111-12 1272

1 gaatteggettteacetgeg......acaceaaaagtaaagaatte 1272

IDENTITY_NUC Gapop 10.0 , Gapext 1.0

Scoring table:

5998275 seqs, 2833880070 residues Searched:

11996550 Total number of hits satisfying chosen parameters:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Database :

/EMC_Celerra_SIDS3/ptodata/2/pna/US11_NEW_COMB.seqz/EMC_Celerra_SIDS3/ptodata/2/pna/US11_EMS_COMB.seqz/EMC_Celerra_SIDS3/ptodata/2/pna/US11_NEW_COMB.seqz//EMC_Celerra_SIDS3/ptodata/2/pna/US10_NEW_COMB.seqz/EMC_Celerra_SIDS3/ptodata/2/pna/US60_NEW_COMB.seqz

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Query Score Match Length DB

Run on:

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3, 2006, 02:41:24; Search time 8110 Seconds (without alignments) 8948.163 Million cell updates/sec
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'EMC_Celerra_SIDS3/ptodata/2/pna/PCTUSC_COMB.
'EMC_Celerra_SIDS3/ptodata/2/pna/VS075_COMB.
GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

SUMMARIES